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Figure 1a hAMGEN2520-59 Coding region and amino acid sequence.

Seq ID NO:

				10			20			3	-			40			50			60
1	ΑT	GCG	AGC	CAC	CCC'	TCT	GGC'	rgc'	TCC'	rgc(GGG	TTC	CCT	GTC	CAG	GAA	GAA	GCG	GTT(GGAG
2	М	R	A	${f T}$	Ρ	L	A	Α	Р	A	G	S	L	S	R	K	K	R	L	\mathbf{E}
				70			80			9	0		1	00			110			120
	фф	CGA	TGA		СТТ	AGA	TAC	CGA	GCG'	TCC	CGT	CCA	GAA	ACG.	AGC'	TCG	AAG:	rgg	GCC	CCAG
	L	D	D	N	L	D	Т	E	R	P	V	Q	K	R	A	R	S	G	Ρ	Q
	ч	ע	_	30		ט	140	11	10	15	-	×		60			170	_		180
	aa	יריז רי			ccc	СТС		ئانىڭ	מממ		-	מממ		-	ጥርር	тсс		TCG	тдс	AACT
		R.	AC1 L	P	P	CIG	L	L	P	L	S	P	P	T	A	P	D	R	A	Т
	P	ĸ		90	Р	C	200	ш	r	21		E	_	20	73.	_	230	10	2 1	240
	~~	mam			maa	аша		пат	шаа			m/dm			~~~	~~		707	aca.	CGGG
																GCC P	E E	E E	GGG	G
	A	V	A	T	A	S	R	L	G	P	Y	V	L	L	E	Р		ᆫ	G	_
				50			260	~-~	~ ~ ~	27	-	~ ~		80	m	с т.	290	aam	~mx	300
	CG																			CCCC
	R	A	Y		A	L	H	C	Р	\mathbf{T}	G	${ m T}$	E	Y	T	С	K	V	Y	P
				10			320			33	-		_	40			350			360
	GT	'CCA	GGA	AGC	CCT:	'GGC	CCGT	GCT	GGA	GCC	CTA	.CGC	GCG		GCC	CCC				TGTG
	V	Q	E	A	L	A	V	L	\mathbf{E}	P	Y	Α	R	\mathbf{L}	P	P	H	K	Η	V
				70			380			39			_	00			410			420
	GC	TCG	GCC	CAC	'TGA	.GG7	CCT	GGC	TGG	TAC	CCA	.GCT	CCT	CTA	.CGC	CTT	CTTT	CAC	TCG	GACC
	A	R	P	\mathbf{T}	E	V	L	A	G	${f T}$	Q	\mathbf{L}	L	Y	Α	F	F	${ m T}$	R	\mathbf{T}
			4	30			440			45	0		4	60			470			480
	CA	TGG	GGA	CAT	'GCA	CAC	GCCT	GGT	'GCG	AAG	CCG	CCA	CCG	TAT	'CCC	TGA	AGCC'	TGA	.GGC	TGCC
	Н	G	D	M	H	S	L	V	R	S	R	H	R	I	P	\mathbf{E}	P	E	A	A
			4	90			500			51	0		5	20			530			540
	GT	GCT	'CTT	CCG	CCA	GA:	rggc	CAC	:CGC	CCT	'GGC	:GCA	.CTG	TCA	CCA	.GCI	ACGG	TCT	'GGT	CCTG
	V	L	F	R	0	М	A	\mathbf{T}	A	L	Α	H	C	H	Q	Н	G	L	V	L
				550	~		560			57	0		5	80			590			600
	CC	TGA	ጥርባ	CAA	каст	'GT(СТТ	ጥርተ	СТТ	'CGC	'TGA	.CCG	TGA	GAG	GAZ	AGAA	GCT	GGT	GCTG
	R	D	Τ,	K	L	C	R	F	V	F	A	D	R	E	R	K	K	L	V	L
			_	510		Ū	620	_	•	63			6	40			650			660
	GZ	Z Z D			CCD	СТС		ССТ	ССТ			;GCC	_		ידידר	laar		GGA	CAA	GCAC
	E	N	L	E	D	S	C	V	L	Т	G	P	D	D	S	L	W	D	K	Н
	11	14		570	יב	D	680	٧	ш	69	_	-		00	~	_	710			720
	CC	чете			א ליידי א	עפי		א כיכ	ייייכיא			יכאכ			ממכר	יריםי		CTC	יממפ	CAAG
	A	C	P	JEA. A	Y	V	G	ACC P	E	I	L L	S	S	R.	A	S	Y	S	G	K
	A	C	_		1	V	740		Ľ	75		S		60	A	J	770		J	780
	0.0	77.00		730	nama	7/17/2			ааап			- Carrier			יממי	100			(TT)	
			CGA	ATGI	r.C.T.C	AU	JCCT	النالنال	iCG L	رحور	ひし1	.CII	CAC	$\mathcal{L}_{\mathbf{M}}$. GC 1	יטט. ת	2000	TT	ZZ ZZ	CCCC P
	A	Α			W	5											G		I	
				790			800													840
																				CCTTG
	F	Q		S	E	P									R	G	A		A	
				350			860							80			890		· ~ · · ·	900
	C	CTGC																		GCCA
	Р	Α	G	L	S	A	P	Α	R	C	L	V	Ŕ	C	L	L	R	R	\mathbf{E}	P

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Figure 1b

hAMGEN2520-59 Coding region and amino acid sequence (continued).

Seq ID NO:

950 940 920 930 910 1 GCTGAACGGCTCACAGCCACAGGCATCCTCCTGCACCCCTGGCTGCGACAGGACCCGATG Q D P Μ L L H P W 2 A E R L T A T G I 1020 1010 980 990 1000 970 CCCTTAGCCCCAACCCGATCCCATCTCTGGGAGGCTGCCCAGGTGGTCCCTGATGGACTG V V P D G L H L W E A Q R S A P 1060 1070 1050 1030 1040 V V L GLDEAREE G D R Ε \mathbf{E}

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Figure 2a

2520-59

Genbank BAB15597

50

LRFAS PGPGAGRARDSQRKWRRLRARPLLGPGQGWSWAGI PSSAAAQRAG --GQGWSWAGIPSSAAAQRAG (1) (1) (1)(1)

(1) Patented Seq: NO 1367 Patented SEQ ID NO:1102 JJ503-KS

--MRATPLAAPAGSLSRKKRLELDDNLDT -----MRATPLAAPAGSLSRKKRLELDDNLDT (1) Consensus 2520-59 Genbank BAB15597

PPAGALEALSPGGARAHAERRGEMRATPLAAPAGSLSRKKRLELDDNLDT (51)(1)Patented SEQ ID NO:1102 Patented Seq: NO 1367

PPAGALEALSPGGARAHAERRGEMRATPLAAPAGSLSRKKRLELDDNLDT (20)JJ503-KS

(51)

Consensus

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> (28)(28)2520-59

101) Genbank BAB15597 Patented Seq: NO 1367

(10) (1)Patented SEQ ID NO:1102 JJ503-KS ERPVOKRARSGPOPRLPPCLLPLSPPTAPDRATAVATASRLGPYVLLEPE (101)Consensus

ERPVOKRARSGPOPRLPPCLLPLSPPTAPDRATAVATASRLGPYVLLEPE

ERPVOKRARSGPQPRLPPCLLPLSPPTAPDRATAVATASRLGPYVLLEPE ERPVOKRARSGPOPRLPPCLLPLSPPTAPDRATAVATASRLGPYVLLEPE ERPVQKRARSGPQPRLPPCLLPLSPPTAPDRATAVXTXSRXXXYVLLEAR

MRATPLAAPAGSLSRKKRLELDDNLDT

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Figure 2b

		151 200
2520-59	(78)	EGGRAYRALHCPTGTEYTCKVYPVQEALAVLEPYARLPPHKHVARPTEVL
Genbank BAB15597	(78)	EGGRAYQALHCPTGTEYTCKVYPVQEAPAVLEPYARLPPHKHVARPTEVL
Patented Seq: NO 1367	(151)	RXA
Patented SEQ ID NO:1102	(1)	d9
JJ503-KS	(120)	EGGRAYQALHCPTGTEYTCKVYPVQEALAVLEPYARLPPHKHVARPTEVL
Consensus	(151)	EGGRAYQALHCPTGTEYTCKVYPVQEA AVLEPYARLPPHKHVARPTEVL
		201 250
2520-59	(128)	AGTQLLYAFFTRTHGDMHSLVRSRHRIPEPEAAVLFRQMATALAHCHQHG
Genbank BAB15597	(128)	AGTQLLYAFFTRTHGDMHSLVRSRHRIPEPEAAVLFRQMATALAHCHQHG
Patented Seq: NO 1367	(154)	
Patented SEQ ID NO:1102	(3)	GWYPAPLRLFHSDPWGHAQPGAKRHRIPEPEAAVLFRQMATALAHCHQHG
JJ503-KS	(170)	AGTQLLYAFFTRTHGDMHSLVRSRHRIPEPEAAVLFRQMATALAHCHQHG
Consensus	(201)	AGTQLLYAFFTRTHGDMHSLVRSRHRIPEPEAAVLFRQMATALAHCHQHG
		251 300
2520-59	(178)	LVLRDLKLCRFVFADRERKKLVLENLEDSCVLTGPDDSLWDKHACPAYVG
Genbank BAB15597	(178)	LVLRDLKLCRFVFADRERKKLVLENLEDSCVLTGPDDSLWDKHACPAYVG
Patented Seq: NO 1367	(154)	
Patented SEQ ID NO:1102	(53)	LVLRDLKLCRFVFADRERKKLVLENLEDSCVLTGPDDSLWDKHACPAYVG
JJ503-KS	(220)	LVLRDLKLCRFVFADRERKKLVLENLEDSCVLTGPDDSLWDKHACPAYVG
Consensus	(251)	LVLRDLKLCRFVFADRERKKLVLENLEDSCVLTGPDDSLWDKHACPAYVG
		301 350
2520-59	(228)	PEILSSRASYSGKAADVWSLGVALFTMLAGHYPFQDSEPVLLFGKIRRGA
Genbank BAB15597	(228)	PEILSSRASYSGKAADVWSLGVALFTMLAGHYPFQDSEPVLLFGKIRRGA
Patented Seq: NO 1367	(154)	
Patented SEQ ID NO:1102	(103)	PEILSSRASYSGKAADVWSLGVALFTMLAGHYPFQDSEPVLLFGKIRRGA
JJ503-KS	(270)	PEILSSRASYSGKAADVWSLGVALFTMLAGHYPFQDSEPVLLFGKIRRGA
Consensus	(301)	PEILSSRASYSGKAADVWSLGVALFTMLAGHYPFQDSEPVLLFGKIRRGA

Figure 2c

Title: Novel Serine Threonine Kinase Member, h2520-59 Docket No. 01017/36524A Page 5 of 5 Fig. 2C

400 (278) YALPAGLSAPARCLVRCLLRREPAERLTATGILLHPWLRQDPMPLAPTRS (278) YALPAGLSAPARCLVRCLLRREPAERLTATGILLHPWLRQDPMPLAPTRS (154)	(328) HLWEAAQVVPDGLGLDEAREEEGDREVVLYG (328) HLWEAAQVVPDGLGLDEAREEEGDREVVLYG (154)
351 YALPAGLS YALPAGLS YALPAGLS YALPAGLS YALPAGLS	HLWEAAQVVPI HLWEAAQVVPI HLWEAAQVVPI HLWEAAQVVPI
(278)	(328)
(278)	(328)
(154)	(154)
(153)	(203)
(320)	(361)
(351)	(401)
2520-59	2520-59
Genbank BAB15597	Genbank BAB15597
Patented Seq: NO 1367	Patented Seq: NO 1367
Patented SEQ ID NO:1102	Patented SEQ ID NO:1102
JJ503-KS	JJ503-KS

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